TECH CENTER 1600/2900

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<110> Estell, David A.
Harding, Fiona A.
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<120> PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND METHODS OF MAKING AND USING THE SAME

<130> A-68893/DJB/DAV

<140> US 09/500,135

<141> 2000-02-08

<150> US 09/060,872

<151> 1998-04-15

<160> 236

<170> PatentIn Ver. 2.1

<210> 1

<211> 1495

<212> DNA

<213> Bacillus amyloliquefaciens

<220>

<221> mat_peptide

<222> (417)..(1495)

<220>

<221> CDS

<222> (96)..(1244)

<220>

<221> misc feature

<222> (582)..(584)

<223> The nnn at positions 582 through 584 which in a preferred embodiment (aat) is to code for asparagine, but which may also code for proline.

<220>

<221> misc feature

<222> (585)..(587)

<223> The nnn at positions 585 through 587 which in a
 preferred embodiment (cct) is to code for proline,
 but which may also code for asparagine.

<220>

<221> misc feature

<222> (597)..(599)

<223> The nnn at positions 597 to 599 which in a
 preferred embodiment (aac) is to code for
 asparagine, but which may also code for aspartic acid.

<220>

<221> misc feature

<222> (678)..(680)

<223> The nnn at positions 678 through 680 which in a preferred embodiment (gca) is to code for alanine, but which may also code for serine. <220> <221> misc feature <222> (681)..(683) <223> The nnn at positions 681 through 683 which in a preferred embodiment (tca) is to code for serine, but which may also code for alanine. <220> <221> misc_feature <222> (708)..(710) <223> The nnn at positions 708 through 710 which in a preferred embodiment (gct) is to code for alanine, but which may also code for aspartic acid. <220> <221> misc feature <222> (711) .. (713) <223> The nnn at positions 711 through 713 which in a preferred embodiment (gac) is to code for aspartic acid, but which may also code for alanine. <220> <221> misc_feature <222> (888)..(890) <223> The nnn at positions 888 through 890 which in a preferred embodiment (act) is to code for threonine, but which may also code for serine. <220> <221> misc feature <222> (891)..(893) <223> The nnn at positions 891 through 893 which in a preferred embodiment (tcc) is to code for serine, but which may also code for threonine. <220> <221> misc feature <222> (1167)..(1169) <223> The nnn at positions 1167 through 1169 which in a preferred embodiment (gaa) is to code for

<400> 1

ggtctactaa aatattattc catactatac aattaataca cagaataatc tgtctattgg 60

glutamic acid, but which may also code for glutamine.

ttattctgca aatgaaaaaa aggagaggat aaaga atg aga ggc aaa aaa gta 113 Met Arg Gly Lys Lys Val -105

tgg atc agt ttg ctq ttt gct tta gcg tta atc ttt acg atg gcg ttc 161
Trp Ile Ser Leu Leu Phe Ala Leu Ala Leu Ile Phe Thr Met Ala Phe
-100 -95 -90

				tct Ser												209
				ggg Gly -65												257
				gtc Val												305
				gac Asp												353
				aaa Lys												401
				tac Tyr -1												449
				ctg Leu												497
				gac Asp												545
				gcc Ala												593
gac Asp 60	nnn Xaa	aac Asn	tct Ser	cac His	gga Gly 65	act Thr	cac His	gtt Val	gcc Ala	ggc Gly 70	aca Thr	gtt Val	gcg Ala	gct Ala	ctt Leu 75	641
				ggt Gly 80												689
				ctc Leu												737
att Ile	aac Asn	gga Gly 110	atc Ile	gag Glu	tgg Trp	gcg Ala	atc Ile 115	gca Ala	aac Asn	aat Asn	atg Met	gac Asp 120	gtt Val	att Ile	aac Asn	785
				gga Gly												833
gat	aaa	gcc	gtt	gca	tcc	ggc	gtc	gta	gtc	gtt	gcg	gca	gcc	ggt	aac	881

Asp Lys Ala Val Ala Ser Gly Val Val Val Val Ala Ala Ala Gly Asn 140 145 150 155													
gaa ggc nnn nnn ggc agc tca agc aca gtg ggc tac cct ggt aaa tac 929 Glu Gly Xaa Xaa Gly Ser Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr 160 165 170													
cct tct gtc att gca gta ggc gct gtt gac agc agc aac caa aga gca 977 Pro Ser Val Ile Ala Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala 175 180 185													
tct ttc tca agc gta gga cct gag ctt gat gtc atg gca cct ggc gta 102 Ser Phe Ser Ser Val Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val 190 195 200	5												
tct atc caa agc acg ctt cct gga aac aaa tac ggg gcg tac aac ggt 107 Ser Ile Gln Ser Thr Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly 205 210 215	3												
acg tca atg gca tct ccg cac gtt gcc gga gcg gct gct ttg att ctt 112 Thr Ser Met Ala Ser Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu 220 235 230 235	1												
tct aag cac ccg aac tgg aca aac act caa gtc cgc agc agt tta nnn 116 Ser Lys His Pro Asn Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Xaa 240 245 250	9												
aac acc act aca aaa ctt ggt gat tct ttc tac tat gga aaa ggg ctg 121 Asn Thr Thr Thr Lys Leu Gly Asp Ser Phe Tyr Tyr Gly Lys Gly Leu 255 260 265	7												
atc aac gta cag gcg gca gct cag taa aacataaaaa accggccttg 126 Ile Asn Val Gln Ala Ala Gln 270 275	4												
gccccgccgg tttttttatt tttcttcctc cgcatgttca atccgctcca taatcgacgg 132	4												
atggeteect etgaaaattt taaegagaaa eggegggttg aeeeggetea gteeegtaae 1													
ggccaagtcc tgaaacgtct caatcgccgc ttcccggttt ccggtcagct caatgccgta													
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<223> Xaa = Asn or Pro <220> <221> VARIANT <222> (164)(164)													

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<221> VARIANT
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<221> VARIANT
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<223> Xaa = Ala or Ser
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<221> VARIANT
<222> (196)...(196)
<223> Maa = Ser or Ala
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<221> VAP.IANT
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<222> (265)...(265)
<223> Maa = Thr or Ser
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Ile Phe Thr Met Ala Phe Gly Ser Thr Ser Ser Ala Gln Ala Ala Gly
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Lys Ser Asn Gly Glu Lys Lys Tyr Ile Val Gly Phe Lys Gln Thr Met
                              40
Ser Thr Met Ser Ala Ala Lys Lys Lys Asp Val Ile Ser Glu Lys Gly
                          55
Gly Lys Val Gln Lys Gln Phe Lys Tyr Val Asp Ala Ala Ser Ala Thr
                                           75
                     70
Leu Asn Glu Lys Ala Val Lys Glu Leu Lys Lys Asp Pro Ser Val Ala
                                      90
Tyr Val Glu Glu Asp His Val Ala His Ala Tyr Ala Gln Ser Val Pro
```

105

Tyr Gly Val Ser Gln Ile Lys Ala Pro Ala Leu His Ser Gln Gly Tyr 120 Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp Ser Gly Ile Asp Ser 135 Ser His Pro Asp Leu Lys Val Ala Gly Gly Ala Ser Met Val Pro Ser 155 150 Glu Thr Xaa Xaa Phe Gln Asp Xaa Asn Ser His Gly Thr His Val Ala 165 170 Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val Ala 185 180 Pro Ser Xaa Xaa Leu Tyr Ala Val Lys Val Leu Gly Xaa Xaa Gly Ser 200 Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu Trp Ala Ile Ala Asn 215 220 Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly Pro Ser Gly Ser Ala 235 230 Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala Ser Gly Val Val Val 250 255 245 Val Ala Ala Gly Asn Glu Gly Xaa Xaa Gly Ser Ser Ser Thr Val 270 265 Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala Val Gly Ala Val Asp 280 Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val Gly Pro Glu Leu Asp 300 295 Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr Leu Pro Gly Asn Lys 310 315 Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser Pro His Val Ala Gly 330 325 Ala Ala Leu Ile Leu Ser Lys His Pro Asn Trp Thr Asn Thr Gln 340 345 350 Val Arg Ser Ser Leu Xaa Asn Thr Thr Thr Lys Leu Gly Asp Ser Phe 355 360 Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala Ala Ala Gln 375

<210> 3 <211> 275 <212> PRT

<213> Bacillus amyloliquefaciens

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Ala Gln Ser Val Pro Tyr Gly Val Ser Gln Ile Lys Ala Pro Ala Leu 1 5 10 15

His Ser Gln Gly Tyr Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp 20 25 30

Ser Gly Ile Asp Ser Ser His Pro Asp Leu Lys Val Ala Gly Gly Ala 35 40 45

Ser Met Val Pro Ser Glu Thr Asn Pro Phe Gln Asp Asn Asn Ser His
50 55 60

Gly Thr His Val Ala Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly 65 70 75 80 Val Leu Gly Val Ala Pro Ser Ala Ser Leu Tyr Ala Val Lys Val Leu 85 90 95

Gly Ala Asp Gly Ser Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu 100 105 110

Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly
115 120 125

Pro Ser Gly Ser Ala Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala 130 135 140

Ser Gly Val Val Val Val Ala Ala Ala Gly Asn Glu Gly Thr Ser Gly 145 150 155 160

Ser Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala 165 170 175

Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val 180 185 190

Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr 195 200 205

Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser 210 215 220

Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn 225 230 235 240

Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Glu Asn Thr Thr Thr Lys 245 250 255

Leu Gly Asp Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala
260 265 270

Ala Ala Gln 275

<210> 4

<211> 275

<212> PRT

<213> Bacillus subtilis

<400> 4

Ala Gln Ser Val Pro Tyr Gly Ile Ser Gln Ile Lys Ala Pro Ala Leu 1 5 10 15

His Ser Gln Gly Tyr Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp 20 25 30

Ser Gly Ile Asp Ser Ser His Pro Asp Leu Asn Val Arg Gly Gly Ala

Ser Phe Val Pro Ser Glu Thr Asn Pro Tyr Gln Asp Gly Ser Ser His

50

Gly 65	Thr	His	Val	Ala	Gly 70	Thr	Ile	Ala	Ala	Leu 75	Asn	Asn	Ser	Ile	Gly 80
Val	Leu	Gly	Val	Ser 85	Pro	Ser	Ala	Ser	Leu 90	Tyr	Ala	Val	Lys	Val 95	Leu
Asp	Ser	Thr	Gly 100	Ser	Gly	Gln	Tyr	Ser 105	Trp	Ile	Ile	Asn	Gly 110	Ile	Glu
Trp	Ala	Ile 115	Ser	Asn	Asn	Met	Asp 120	Val	Ile	Asn	Met	Ser 125	Leu	Gly	Gly
Pro	Thr 130	Gly	Ser	Thr	Ala	Leu 135	Lys	Thr	Val	Val	Asp 140	Lys	Ala	Val	Ser
Ser 145	Gly	Ile	Val	Val	Ala 150	Ala	Ala	Ala	Gly	Asn 155	Glu	Gly	Ser	Ser	Gly 160
Ser	Thr	Ser	Thr	Val 165	Gly	Tyr	Pro	Ala	Lys 170	Tyr	Pro	Ser	Thr	Ile 175	Ala
Val	Gly	Ala	Val 180	Asn	Ser	Ser	Asn	Gln 185	Arg	Ala	Ser	Phe	Ser 190	Ser	Ala
Gly	Ser	Glu 195	Leu	Asp	Val	Met	Ala 200	Pro	Gly	Val	Ser	Ile 205	Gln	Ser	Thr
Leu	Pro 210	Gly	Gly	Thr	Tyr	Gly 215	Ala	Tyr	Asn	Gly	Thr 220	Ser	Met	Ala	Thr
Pro 225	His	Val	Ala	Gly	Ala 230	Ala	Ala	Leu	Ile	Leu 235	Ser	Lys	His	Pro	Thr 240
Trp	Thr	Asn	Ala	Gln 245	Val	Arg	Asp	Arg	Leu 250	Glu	Ser	Thr	Ala	Thr 255	Tyr
Leu	Gly	Asn	Ser 260	Phe	Tyr	Tyr	Gly	Lys 265	Gly	Leu	Ile	Asn	Val 270	Gln	Ala
Ala	Ala	Gln 275													
<210> 5 <211> 274 <212> PRT <213> Bacillus licheniformis															
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55

60

Gln Ala Gln Gly Phe Lys Gly Ala Asn Val Lys Val Ala Val Leu Asp 20 25 30 Thr Gly Ile Gln Ala Ser His Pro Asp Leu Asn Val Val Gly Gly Ala 35 40 45

Ser Phe Val Ala Gly Glu Ala Tyr Asn Thr Asp Gly Asn Gly His Gly 50 55 60

Thr His Val Ala Gl; Thr Val Ala Ala Leu Asp Asn Thr Thr Gly Val 65 70 75 80

Leu Gly Val Ala Pro Ser Val Ser Leu Tyr Ala Val Lys Val Leu Asn 85 90 95

Ser Ser Gly Ser Gl; Ser Tyr Ser Gly Ile Val Ser Gly Ile Glu Trp 100 105 110

Ala Thr Thr Asn Gly Met Asp Val Ile Asn Met Ser Leu Gly Gly Ala 115 120 125

Ser Gly Ser Thr Ala Met Lys Gln Ala Val Asp Asn Ala Tyr Ala Arg 130 135 140

Gly Val Val Val Ala Ala Ala Gly Asn Ser Gly Asn Ser Gly Ser 145 150 155 160

Thr Asn Thr Ile Gly Tyr Pro Ala Lys Tyr Asp Ser Val Ile Ala Val 165 170 175

Gly Ala Val Asp Ser Asn Ser Asn Arg Ala Ser Phe Ser Ser Val Gly
180 185 190

Ala Glu Leu Glu Val Met Ala Pro Gly Ala Gly Val Tyr Ser Thr Tyr 195 200 205

Pro Thr Asn Thr Tyr Ala Thr Leu Asn Gly Thr Ser Met Ala Ser Pro 210 215 220

His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn Leu 225 230 235 240

Ser Ala Ser Gln Val Arg Asn Arg Leu Ser Ser Thr Ala Thr Tyr Leu 245 250 255

Gly Ser Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Glu Ala Ala 260 265 270

Ala Gln

<210> 6

<211> 269

<212> PRT

<213> Human

<400> 6

Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala 1 5 10 15 His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp 20 25 30

Thr Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser 35 40 45

Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly Thr 50 55 60

His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu 65 70 75 80

Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala 85 90 95

Ser Gly Ser Gly Ser Val Ser Ser Ile Ala Gln Gly Leu Glu Trp Ala 100 105 110

Gly Asn Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser Pro Ser 115 120 125

Pro Ser Ala Thr Leu Glu Gln Ala Val Asn Ser Ala Thr Ser Arg Gly 130 135 140

Val Leu Val Val Ala Ala Ser Gly Asn Ser Gly Ala Gly Ser Ile Ser 145 150 155 160

Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly Ala Thr Asp Gln
165 170 175

Asn Asn Asn Arg Ala Ser Phe Ser Gln Tyr Gly Ala Gly Leu Asp Ile 180 185 190

Val Ala Pro Gly Val Asn Val Gln Ser Thr Tyr Pro Gly Ser Thr Tyr 195 200 205

Ala Ser Leu Asn Gly Thr Ser Met Ala Thr Pro His Val Ala Gly Ala 210 215 220

Ala Ala Leu Val Lys Gln Lys Asn Pro Ser Trp Ser Asn Val Gln Ile 225 230 235 240

Arg Asn His Leu Lys Asn Thr Ala Thr Ser Leu Gly Ser Thr Asn Leu 245 250 255

Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala Thr Arg 260 265

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<211> 15

<212> PRT

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<220>

<223> Description of Artificial Sequence: Synthetic

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Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala
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Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala His Asn
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Gly Ile Ser Arg Val Gln Ala Pro Ala Ala His Asn Arg Gly Leu
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Arg Val Gln Ala Pro Ala Ala His Asn Arg Gly Leu Thr Gly Ser
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        5
<210> 13
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Ala Pro Ala Ala His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys
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<210> 14
<211> 15
<212> PRT
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Ala His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val
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<210> 15
<211> 15
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Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp Thr Gly Ile Ser
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Asp Leu Asn Ile Arg Gly Gly Ala Ser Phe Val Pro Gly Glu Pro
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Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly
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1 5
<210> 27
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Ser Thr Gln Asp Gly Asn Gly His Gly Thr His Val Ala Gly Thr
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<211> 15
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Asp Gly Asn Gly His Gly Thr His Val Ala Gly Thr Ile Ala Ala
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<220>

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Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val Ala Pro
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<211> 15
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Leu Asn Asn Ser Ile Gly Val Leu Gly Val Ala Pro Ser Ala Glu
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Val Leu Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val
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<210> 36
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<212> PRT
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<211> 15
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Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser
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Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser Gly Ser Val
1 5
                                   10
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Ser Ser Ile Ala Gln Gly Leu Glu Trp Ala Gly Asn Asn Gly Met
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          5
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<211> 15
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Ala Gln Gly Leu Glu Trp Ala Gly Asn Asn Gly Met His Val Ala
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<210> 45
<211> 15
<212> PRT
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Leu Glu Trp Ala Gly Asn Asn Gly Met His Val Ala Asn Leu Ser
1
                 5
<210> 46
<211> 15
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Ala Gly Asn Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser
            5
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.

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<210> 47
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Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser Pro Ser Pro
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<211> 15
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His Val Ala Asn Leu Ser Leu Gly Ser Pro Ser Pro Ser Ala Thr
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<210> 49
<211> 15
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Val Gln Ser Thr Tyr Pro Gly Ser Thr Tyr Ala Ser Leu Asn Gly
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Val Gln Ile Arg Asn His Leu Lys Asn Thr Ala Thr Ser Leu Gly
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Ser Thr Asn Leu Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala
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Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn Asn Gln Val
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Pro Leu Arg Arg Ala Ser Leu Ser Leu Gly Ser Gly Phe Trp His
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Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu Trp Gln Met
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Leu Gln Ala Asp Val Leu Trp Gln Met Gly Tyr Thr Gly Ala Asn
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Asp Val Leu Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val
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His Phe Lys Asn Val Lys Glu Arg Thr Asn Trp Thr Asn Glu Arg
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Glu Arg Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp Asp Gly Leu
                                     10
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<223> Description of Artificial Sequence: Synthetic
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                                    10
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Thr Leu Asp Asp Gly Leu Gly His Gly Thr Phe Val Ala Gly Val
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                                     10
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<212> PRT
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Asp Gly Leu Gly His Gly Thr Phe Val Ala Gly Val Ile Ala Ser
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<210> 125
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Gly His Gly Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu
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Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly
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Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly Phe Ala Pro
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Ile Ala Ser Met Arg Glu Cys Gln Gly Phe Ala Pro Asp Ala Glu
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Met Arg Glu Cys Gln Gly Phe Ala Pro Asp Ala Glu Leu His Ile
                                   10
<210> 130
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Asn Gln Val Ser Tyr Thr Ser Trp Phe Leu Asp Ala Phe Asn Tyr
 1 5
                         10
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Ser Trp Phe Leu Asp Ala Phe Asn Tyr Ala Ile Leu Lys Lys Ile
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Asn Leu Ser Ile Gly Gly Pro Asp Phe Met Asp His Pro Phe Val
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Ile Gly Gly Pro Asp Phe Met Asp His Pro Phe Val Asp Lys Val
        5
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                                   10
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Met Asp His Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn
                 5
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Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile
              5
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Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile Met Val Ser
                5
                                   10
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Trp Glu Leu Thr Ala Asn Asn Val Ile Met Val Ser Ala Ile Gly
                                    10
                                                        15
                5
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Thr Ala Asn Asn Val Ile Met Val Ser Ala Ile Gly Asn Asp Gly
          5
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Asn Val Ile Met Val Ser Ala Ile Gly Asn Asp Gly Pro Leu Tyr
 1 5
                                   10
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<400> 153
Met Val Ser Ala Ile Gly Asn Asp Gly Pro Leu Tyr Gly Thr Ile
                                    10
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                 5
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Ala Ile Gly Asn Asp Gly Pro Leu Tyr Gly Thr Leu Asn Asn Pro
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Pro Leu Tyr Gly Thr Leu Asn Asn Pro Ala Asp Gln Met Asp Val
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<210> 157
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Gly Thr Leu Asn Asn Pro Ala Asp Gln Met Asp Val Ile Gly Val
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<210> 159
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<223> Description of Artificial Sequence: Synthetic
<400> 159
Ala Asp Gln Met Asp Val Ile Gly Val Gly Gly Ile Asp Phe Glu
<210> 160
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<212> PRT
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Met Asp Val Ile Gly Val Gly Gly Ile Asp Phe Glu Asp Asn Ile
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USSN 09/500,135
<210> 161
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<211> 15

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic

<400> 161

Ile Gly Val Gly Gly Ile Asp Phe Glu Asp Asn Ile Ala Arg Phe 10

<210> 162

<211> 15

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Synthetic

<400> 162

Gly Gly Ile Asp Phe Glu Asp Asn Ile Ala Arg Phe Ser Ser Arg 10

<210> 163

<211> 15

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<213> Artificial Sequence

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<400> 163

Asp Phe Glu Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr 5

<210> 164

<211> 15

<212> PRT

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<223> Description of Artificial Sequence: Synthetic

<400> 164

Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr Thr Trp Glu 1 5 10

<210> 165

<211> 15

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USSN 09/500,135

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Ser Ser Arg Gly Met Thr Trp Glu Leu Pro Gly Gly Tyr Gly
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Gly Met Thr Trp Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys
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<400> 168
Thr Trp Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile
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Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile Val Thr Tyr
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<400> 174
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<210> 175
<211> 15
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Ser Gly Val Lys Gly Gly Cys Arg Ala Leu Ser Gly Thr Ser Val
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<400> 177
Lys Gly Gly Cys Arg Ala Leu Ser Gly Thr Ser Val Ala Ser Pro
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Cys Arg Ala Leu Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala
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Leu Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala Gly Ala Val
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Thr Ser Val Ala Ser Pro Val Val Ala Gly Ala Val Thr Leu Leu
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<210> 181
<211> 15
<212> PRT
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<400> 181
Ala Ser Pro Val Val Ala Gly Ala Val Thr Leu Leu Val Ser Thr
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<210> 182
<211> 15
<212> PRT
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Val Val Ala Gly Ala Val Thr Leu Leu Val Ser Thr Val Gln Lys
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Gly Ala Val Thr Leu Leu Val Ser Thr Val Gln Lys Arg Glu Leu
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Thr Leu Leu Val Ser Thr Val Gln Lys Arg Glu Leu Val Asn Pro
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Val Ser Thr Val Gln Lys Arg Glu Leu Val Asn Pro Ala Ser Met
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Val Gln Lys Arq Glu Leu Val Asn Pro Ala Ser Met Lys Gln Ala
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<223> Description of Artificial Sequence: Synthetic
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Arg Glu Leu Val Asn Pro Ala Ser Met Lys Gln Ala Leu Ile Ala
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Val Asn Pro Ala Ser Met Lys Gln Ala Leu Ile Ala Ser Ala Arg
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Ala Ser Met Lys Gln Ala Leu Ile Ala Ser Ala Arg Arg Leu Pro
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<211> 15
<212> PRT
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<400> 190
Lys Gln Ala Leu Ile Ala Ser Ala Arg Arg Leu Pro Gly Val Asn
<210> 191
<211> 15
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<210> 196

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- <400> 196
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- <210> 197
- <211> 15
- <212> PRT
- <213> Artificial Sequence
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- Gly Lys Leu Asp Leu Leu Arg Ala Tyr Gln Ile Leu Asn Ser Tyr 1 5 10 15
- <210> 198
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- <223> Description of Artificial Sequence: Synthetic
- <400> 198
- Asp Leu Leu Arg Ala Tyr Gln Ile Leu Asn Ser Tyr Lys Pro Gln
 1 5 10 15
- <210> 199
- <211> 15
- <212> PRT
- <213> Artificial Sequence
- <220>
- <223> Description of Artificial Sequence: Synthetic
- <400> 199
- Arg Ala Tyr Gln Ile Leu Asn Ser Tyr Lys Pro Gln Ala Ser Leu 1 5 10 15
- <210> 200
- <211> 15
- <212> PRT
- <213> Artificial Sequence
- <220>

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<400> 201
Asn Ser Tyr Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp
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<210> 202
<211> 15
<212> PRT
<213> Artificial Sequence
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Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu
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<210> 203
<211> 15
<212> PRT
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<400> 203
Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr
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<210> 204
<211> 15
<212> PRT
<213> Artificial Sequence
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Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro
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<223> Description of Artificial Sequence: Synthetic
<400> 205
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<210> 206
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<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
<400> 206
Leu Thr Glu Cys Pro Tyr Met Trp Pro Tyr Cys Ser Gln Pro Ile
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<210> 207
<211> 15
<212> PRT
<213> Artificial Sequence
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<210> 208
<211> 1052
<212> PRT
<213> Homo sapiens
<400> 208
Met Lys Leu Val Asn Ile Trp Leu Leu Leu Val Val Leu Leu Cys
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Gly Lys Lys His Leu Gly Asp Arg Leu Glu Lys Lys Ser Phe Glu Lys
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Ala Pro Cys Pro Gly Cys Ser His Leu Thr Leu Lys Val Glu Phe Ser
        35 40
Ser Thr Val Val Glu Tyr Glu Tyr Ile Val Ala Phe Asn Gly Tyr Phe
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Τ	hr 65	Ala	Lys	Ala	Arg	Asn 70	Ser	Phe	Ile	Ser	Ser 75	Ala	Leu	Lys	Ser	Ser 80
0	Slu	Val	Asp	Asn	Trp 85	Arg	Ile	Ile	Pro	Arg 90	Asn	Asn	Pro	Ser	Ser 95	Asp
Γ	yr	Pro	Ser	Asp 100	Phe	Glu	Val	Ile	Gln 105	Ile	Lys	Glu	Lys	Gln 110	Lys	Ala
C	Sly	Leu	Leu 115	Thr	Leu	Glu	Asp	His 120	Pro	Asn	Ile	Lys	Arg 125	Val	Thr	Pro
G	Sln	Arg 130	Lys	Val	Phe	Arg	Ser 135	Leu	Lys	Tyr	Ala	Glu 140	Ser	Asp	Pro	Thr
	/al .45	Pro	Cys	Asn	Glu	Thr 150	Arg	Trp	Ser	Gln	Lys 155	Trp	Gln	Ser	Ser	Arg 160
F	ro	Leu	Arg	Arg	Ala 165	Ser	Leu	Ser	Leu	Gly 170	Ser	Gly	Phe	Trp	His 175	Ala
Τ	hr	Gly	Arg	His 180	Ser	Ser	Arg	Arg	Leu 185	Leu	Arg	Ala	Ile	Pro 190	Arg	Glr
V	al,	Ala	Gln 195	Thr	Leu	Gln	Ala	Asp 200	Val	Leu	Trp	Gln	Met 205	Gly	Tyr	Thr
0	Sly	Ala 210	Asn	Val	Arg	Val	Ala 215	Val	Phe	Asp	Thr	Gly 220	Leu	Ser	Glu	Lys
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Ι	le	Ala	Ser	Met 260	Arg	Glu	Cys	Gln	Gly 265	Phe	Ala	Pro	Asp	Ala 270	Glu	Leu
Н	lis	Ile	Phe 275	Arg	Val	Phe	Thr	Asn 280	Asn	Gln	Val	Ser	Tyr 285	Thr	Ser	Trp
F	he	Leu 290	Asp	Ala	Phe	Asn	Tyr 295	Ala	Ile	Leu	Lys	Lys 300	Ile	Asp	Val	Leu
	sn 05	Leu	Ser	Ile	Gly	Gly 310	Pro	Asp	Phe	Met	Asp 315	His	Pro	Phe	Val	Asp 320
I	ys	Val	Trp	Glu	Leu 325	Thr	Ala	Asn	Asn	Val 330	Ile	Met	Val	Ser	Ala 335	Ile
C	ly	Asn	Asp	Gly 340	Pro	Leu	Tyr	Gly	Thr 345	Leu	Asn	Asn	Pro	Ala 350	Asp	Gln
ľ	let	Asp	Val	Ile	Gly	Val	Gly	Gly	Ile	Asp	Phe	Glu	Asp	Asn	Ile	Ala

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Gly 385	Arg	Met	Lys	Pro	Asp 390	Ile	Val	Thr	Tyr	Gly 395	Ala	Gly	Val	Arg	Gl ₃
Ser	Gly	Val	Lys	Gly 405	Gly	Cys	Arg	Ala	Leu 410	Ser	Gly	Thr	Ser	Val 415	Ala
Ser	Pro	Val	Val 420	Ala	Gly	Ala	Val	Thr 425	Leu	Leu	Val	Ser	Thr 430	Val	Glr
Lys	Arg	Glu 435	Leu	Val	Asn	Pro	Ala 440	Ser	Met	Lys	Gln	Ala 445	Leu	Ile	Ala
Ser	Ala 450	Arg	Arg	Leu	Pro	Gly 455	Val	Asn	Met	Phe	Glu 460	Gln	Gly	His	Gly
Lys 465	Leu	Asp	Leu	Leu	Arg 470	Ala	Tyr	Gln	Ile	Leu 475	Asn	Ser	Tyr	Lys	Pro 480
Gln	Ala	Ser	Leu	Ser 485	Pro	Ser	Tyr	Ile	Asp 490	Leu	Thr	Glu	Cys	Pro 495	Туз
Met	Trp	Pro	Tyr 500	Cys	Ser	Gln	Pro	Ile 505	Tyr	Tyr	Gly	Gly	Met 510	Pro	Thi
Val	Val	Asn 515	Val	Thr	Ile	Leu	Asn 520	Gly	Met	Gly	Val	Thr 525	Gly	Arg	Ile
Val	Asp 530	Lys	Pro	Asp	Trp	Gln 535	Pro	Tyr	Leu	Pro	Gln 540	Asn	Gly	Asp	Asr
Ile 545	Glu	Val	Ala	Phe	Ser 550	Tyr	Ser	Ser	Val	Leu 555	Trp	Pro	Trp	Ser	Gl ₃ 560
Tyr	Leu	Ala	Ile	Ser 565	Ile	Ser	Val	Thr	Lys 570	Lys	Ala	Ala	Ser	Trp 575	Glu
Gly	Ile	Ala	Gln 580	Gly	His	Val	Met	Ile 585	Thr	Val	Ala	Ser	Pro 590	Ala	Glu
Thr	Glu	Ser 595	Lys	Asn	Gly	Ala	Glu 600	Gln	Thr	Ser	Thr	Val 605	Lys	Leu	Pro
Ile	Lys 610	Val	Lys	Ile	Ile	Pro 615	Thr	Pro	Pro	Arg	Ser 620	Lys	Arg	Val	Leu
Trp 625	Asp	Gln	Tyr	His	Asn 630	Leu	Arg	Tyr	Pro	Pro 635	Gly	Tyr	Phe	Pro	Arg 640
Asp	Asn	Leu	Arg	Met 645	Lys	Asn	Asp	Pro	Leu 650	Asp	Trp	Asn	Gly	Asp 655	His
Ile	His	Thr	Asn	Phe	Arq	Asp	Met	Tyr	Gln	His	Leu	Arg	Ser	Met	Gly

660 665 670 Tyr Phe Val Glu Val Leu Gly Ala Pro Phe Thr Cys Phe Asp Ala Ser 675 680 Gln Tyr Gly Thr Leu Leu Met Val Asp Ser Glu Glu Glu Tyr Phe Pro 695 Glu Glu Ile Ala Lys Leu Arg Arg Asp Val Asp Asn Gly Leu Ser Leu Val Ile Phe Ser Asp Trp Tyr Asn Thr Ser Val Met Arg Lys Val Lys 730 725 Phe Tyr Asp Glu Asn Thr Arg Gln Trp Trp Met Pro Asp Thr Gly Gly 745 Ala Asn Ile Pro Ala Leu Asn Glu Leu Leu Ser Val Trp Asn Met Gly Phe Ser Asp Gly Leu Tyr Glu Gly Glu Phe Thr Leu Ala Asn His Asp Met Tyr Tyr Ala Ser Gly Cys Ser Ile Ala Lys Phe Pro Glu Asp Gly 795 Val Val Ile Thr Gln Thr Phe Lys Asp Gln Gly Leu Glu Val Leu Lys Gln Glu Thr Ala Val Val Glu Asn Val Pro Ile Leu Gly Leu Tyr Gln 825 Ile Pro Ala Glu Gly Gly Arg Ile Val Leu Tyr Gly Asp Ser Asn 835 Cys Leu Asp Asp Ser His Arg Gln Lys Asp Cys Phe Trp Leu Leu Asp Ala Leu Leu Gln Tyr Thr Ser Tyr Gly Val Thr Pro Pro Ser Leu Ser 865 870 875 His Ser Gly Asn Arg Gln Arg Pro Pro Ser Gly Ala Gly Ser Val Thr Pro Glu Arg Met Glu Gly Asn His Leu His Arg Tyr Ser Lys Val Leu 905 Glu Ala His Leu Gly Asp Pro Lys Pro Arg Pro Leu Pro Ala Cys Pro 915 Arg Leu Ser Trp Ala Lys Pro Gln Pro Leu Asn Glu Thr Ala Pro Ser 935 Asn Leu Trp Lys His Gln Lys Leu Leu Ser Ile Asp Leu Asp Lys Val 945 950 955 960 Val Leu Pro Asn Phe Arg Ser Asn Arg Pro Gln Val Arg Pro Leu Ser

965 970 975

Pro Gly Glu Ser Gly Ala Trp Asp Ile Pro Gly Gly Ile Met Pro Gly 980 985 990

Arg Tyr Asn Gln Glu Val Gly Gln Thr Ile Pro Val Phe Ala Phe Leu 995 1000 1005

Gly Ala Met Val Val Leu Ala Phe Phe Val Val Gln Ile Asn Lys Ala 1010 1015 1020

Lys Ser Arg Pro Lys Arg Arg Lys Pro Arg Val Lys Arg Pro Gln Leu 1025 1030 1035 1040

Met Gln Gln Val His Pro Pro Lys Thr Pro Ser Val 1045 1050

<210> 209

<211> 280

<212> PRT

<213> Homo sapiens

<400> 209

Arg Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu 1 5 10 15

Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe Asp 20 25 30

Thr Gly Leu Ser Glu Lys His Pro His Phe Lys Asn Val Lys Glu Arg

Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp Asp Gly Leu Gly His Gly 50 55 60

Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly Phe 65 70 75 80

Ala Pro Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn Asn Gln 85 90 95

Val Ser Tyr Thr Ser Trp Phe Leu Asp Ala Phe Asn Tyr Ala Ile Leu 100 105 110

Lys Lys Ile Asp Val Leu Asn Leu Ser Ile Gly Gly Pro Asp Phe Met 115 120 125

Asp His Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val 130 135 140

Ile Met Val Ser Ala Ile Gly Asn Asp Gly Pro Leu Tyr Gly Thr Leu 145 150 155 160

Asn Asn Pro Ala Asp Gln Met Asp Val Ile Gly Val Gly Gly Ile Asp 165 170 175

Phe Glu Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr Thr Trp 180 185 Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile Val Thr Tyr Gly Ala Gly Val Arg Gly Ser Gly Val Lys Gly Gly Cys Arg Ala Leu 210 215 Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala Gly Ala Val Thr Leu 235 Leu Val Ser Thr Val Gln Lys Arg Glu Leu Val Asn Pro Ala Ser Met 250 Lys Gln Ala Leu Ile Ala Ser Ala Arg Arg Leu Pro Gly Val Asn Met 270 265 260 Phe Glu Gln Gly His Gly Lys Leu 275 <210> 210 <211> 15 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic <400> 210 Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val 5 1.0 <210> 211 <211> 15 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic <400> 211 Ala Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val

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<223> Description of Artificial Sequence: Synthetic

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<210> 217

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Gly Ser Ile Ser Tyr Pro Ala Arg Ala Ala Asn Ala Met Ala Val
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Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Ala Ala Met Ala Val
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Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Ala Val
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<223> Description of Artificial Sequence: Synthetic

<400> 221

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<210> 222

<211> 15

<212> PRT

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Pro Gly Gly Val Ala Tyr Ser Cys Ala Asp Gln Thr Pro Trp Ala 1 5 10 15

<210> 223

<211> 15

<212> PRT

<213> Humicola insolens

<400> 223

Cys Gly Trp Ala Lys Lys Ala Pro Val Asn Gln Pro Val Phe Ser
1 5 10 15

<210> 224

<211> 276

<212> PRT

<213> Humicola insolens

<400> 224

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Val Leu Ala Leu Ala Asp Gly Arg Ser Thr Arg Tyr Trp Asp Cys
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Cys Lys Pro Ser Cys Gly Trp Ala Lys Lys Ala Pro Val Asn Gln Pro 35 40 45

Val Phe Ser Cys Asn Ala Asn Phe Gln Arg Ile Thr Asp Phe Asp Ala 50 55 60

Lys Ser Gly Cys Glu Pro Gly Gly Val Ala Tyr Ser Cys Ala Asp Gln 65 70 75 80

Thr Pro Trp Ala Val Asn Asp Asp Phe Ala Leu Gly Phe Ala Ala Thr 85 90 95

Ser Ile Ala Gly Ser Asn Glu Ala Gly Trp Cys Cys Ala Cys Tyr Glu 100 105 110

Leu Thr Phe Thr Ser Gly Pro Val Ala Gly Lys Lys Met Val Val Gln
115 120 125

Ser Thr Ser Thr Gly Gly Asp Leu Gly Ser Asn His Phe Asp Leu Asn 130 135 140

Ile Pro Gly Gly Gly Val Gly Ile Phe Asp Gly Cys Thr Pro Gln Phe 145 150 155 160

Gly Gly Leu Pro Gly Gln Arg Tyr Gly Gly Ile Ser Ser Arg Asn Glu 165 170 175

Cys Asp Arg Phe Pro Asp Ala Leu Lys Pro Gly Cys Tyr Trp Arg Phe 180 185 190

Asp Trp Phe Lys Asn Ala Asp Asn Pro Ser Phe Ser Phe Arg Gln Val 195 200 205

Gln Cys Pro Ala Glu Leu Val Ala Arg Thr Gly Cys Arg Arg Asn Asp 210 215 220

Asp Gly Asn Phe Pro Ala Val Gln Ile Pro Ser Ser Ser Thr Ser Ser 225 230 235 240

Pro Val Asn Gln Pro Thr Ser Thr Ser Thr Thr Ser Thr Ser Thr Thr 245 250 255

Ser Ser Pro Pro Val Gln Pro Thr Thr Pro Ser Gly Cys Thr Ala Glu 260 265 270

Arg Trp Ala Gln 275

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<211> 18

<212> PRT

<213> Thermomyces lanuginosus

<400> 225

Gly Asp Val Thr Gly Phe Leu Ala Leu Asp Asn Thr Asn Lys Leu Ile 1 5 10 15

Val Leu

<210> 226

<211> 15

<212> PRT

<213> Thermomyces lanuginosus

<400> 226

Ser Ile Glu Asn Trp Ile Gly Asn Leu Asn Phe Asp Leu Lys Glu
1 10 15

<210> 227

<211> 291

<212> PRT

<213> Thermomyces lanuginosus

<400> 227

- Met Arg Ser Ser Leu Val Leu Phe Phe Val Ser Ala Trp Thr Ala Leu
 1 5 10 15
- Ala Ser Pro Ile Arg Arg Glu Val Ser Gln Asp Leu Phe Asn Gln Phe 20 25 30
- Asn Leu Phe Ala Gln Tyr Ser Ala Ala Ala Tyr Cys Gly Lys Asn Asn 35 40 45
- Asp Ala Pro Ala Gly Thr Asn Ile Thr Cys Thr Gly Asn Ala Cys Pro 50 60
- Glu Val Glu Lys Ala Asp Ala Thr Phe Leu Tyr Ser Phe Glu Asp Ser
 65 70 75 80
- Gly Val Gly Asp Val Thr Gly Phe Leu Ala Leu Asp Asn Thr Asn Lys
 85 90 95
- Leu Ile Val Leu Ser Phe Arg Gly Ser Arg Ser Ile Glu Asn Trp Ile 100 105 110
- Gly Asn Leu Asn Phe Asp Leu Lys Glu Ile Asn Asp Ile Cys Ser Gly
 115 120 125
- Cys Arg Gly His Asp Gly Phe Thr Ser Ser Trp Arg Ser Val Ala Asp 130 135 140
- Thr Leu Arg Gln Lys Val Glu Asp Ala Val Arg Glu His Pro Asp Tyr 145 150 155 160
- Arg Val Val Phe Thr Gly His Ser Leu Gly Gly Ala Leu Ala Thr Val 165 170 175
- Ala Gly Ala Asp Leu Arg Gly Asn Gly Tyr Asp Ile Asp Val Phe Ser 180 185 190
- Tyr Gly Ala Pro Arg Val Gly Asn Arg Ala Phe Ala Glu Phe Leu Thr
 195 200 205
- Val Gln Thr Gly Gly Thr Leu Tyr Arg Ile Thr His Thr Asn Asp Ile 210 215 220
- Val Pro Arg Leu Pro Pro Arg Glu Phe Gly Tyr Ser His Ser Ser Pro 225 230 235 240
- Glu Tyr Trp Ile Lys Ser Gly Thr Leu Val Pro Val Thr Arg Asn Asp \$245\$ \$250\$ \$255
- Ile Val Lys Ile Glu Gly Ile Asp Ala Thr Gly Gly Asn Asn Gln Pro
 260 265 270
- Asn Ile Pro Asp Ile Pro Ala His Leu Trp Tyr Phe Gly Leu Ile Gly 275 280 285

Thr Cys Leu 290

<210> 228

<211> 15

<212> PRT

<213> Streptomyces plicatus

<400> 228

Ile Lys Val Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly
1 5 10 15

<210> 229

<211> 313

<212> PRT

<213> Streptomyces plicatus

<400> 229

Met Phe Thr Pro Val Arg Arg Arg Val Arg Thr Ala Ala Leu Ala Leu 1 5 10 15

Ser Ala Ala Ala Leu Val Leu Gly Ser Thr Ala Ala Ser Gly Ala
20 25 30

Ser Ala Thr Pro Ser Pro Ala Pro Ala Pro Ala Pro Ala Pro Val Lys 35 40 45

Gln Gly Pro Thr Ser Val Ala Tyr Val Glu Val Asn Asn Asn Ser Met 50 55 60

Leu Asn Val Gly Lys Tyr Thr Leu Ala Asp Gly Gly Asn Ala Phe 65 70 75 80

Asp Val Ala Val Ile Phe Ala Ala Asn Ile Asn Tyr Asp Thr Gly Thr 85 90 95

Lys Thr Ala Tyr Leu His Phe Asn Glu Asn Val Gln Arg Val Leu Asp 100 105 110

Asn Ala Val Thr Gln Ile Arg Pro Leu Gln Gln Gln Gly Ile Lys Val 115 120 125

Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly Phe Ala Asn Phe 130 135 140

Val Ala Lys Tyr Gly Leu Asp Gly Val Asp Phe Asp Asp Glu Tyr Ala 165 170 175

Glu Tyr Gly Asn Asn Gly Thr Ala Gln Pro Asn Asp Ser Ser Phe Val

His Leu Val Thr Ala Leu Arg Ala Asn Met Pro Asp Lys Ile Ile Ser

195 200 205

Leu Tyr Asn Ile Gly Pro Ala Ala Ser Arg Leu Ser Tyr Gly Gly Val 210 215 220

Asp Val Ser Asp Lys Phe Asp Tyr Ala Trp Asn Pro Tyr Tyr Gly Thr 225 230 235 235

Trp Gln Val Pro Gly Ile Ala Leu Pro Lys Ala Gln Leu Ser Pro Ala 245 250 255

Ala Val Glu Ile Gly Arg Thr Ser Arg Ser Thr Val Ala Asp Leu Ala 260 265 270

Arg Arg Thr Val Asp Glu Gly Tyr Gly Val Tyr Leu Thr Tyr Asn Leu 275 280 285

Asp Gly Gly Asp Arg Thr Ala Asp Val Ser Ala Phe Thr Arg Glu Leu 290 295 300

Tyr Gly Ser Glu Ala Val Arg Thr Pro 305 310

• .

<210> 230

<211> 15

<212> PRT

<213> Bacillus amyloliquefaciens

<400> 230

Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val
1 5 10 15

<210> 231

<211> 15

<212> PRT

<213> Bacillus amyloliquefaciens

<400> 231

Asn Gly Ile Glu Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn 1 5 10 15

<210> 232

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 232

Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp Thr Gly Ile Ser
1 5 10 15

<210> 233

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 233

Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser 1 5 10 15

<210> 234

<211> 17

<212> PRT

<213> Bacillus lentus

<400> 234

Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly
1 5 10 15

Ala

<210> 235

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 235

Gly Ala Gly Leu Asp Ile Val Ala Pro Gly Val Asn Val Gln Ser 1 5 10 15

<210> 236

<211> 272

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Hybrid of Bacillus lentus and Bacillus amyloliquefaciens

<400> 236

Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala 1 5 10 15

His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp 20 25 30

Thr Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser 35 40 45

Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly Thr 50 55 60

His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu 65 70 75 80

Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala 85 90 95

Ser Gly Ser Gly Ser Val Ser Ser Ile Ala Gln Gly Leu Glu Trp Ala 105 Gly Asn Asn Gly Met His Val Ile Asn Met Ser Leu Gly Gly Ser Gly Ser Ala Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala Ser Gly Val Val Val Ala Ala Ala Gly Asn Glu Gly Thr Ser Gly Ser Ser Ser 155 Thr Val Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala Val Gly Ala 165 170 Val Asp Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val Gly Pro Glu 185 Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr Leu Pro Gly 195 Asn Lys Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser Pro His Val 215 Ala Gly Ala Ala Leu Ile Leu Ser Lys His Pro Asn Trp Thr Asn 230 235 Thr Gln Val Arg Ser Ser Leu Glu Asn Thr Thr Thr Lys Leu Gly Asp 245 250 Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala Ala Gln

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